M. tuberculosis re Peptide modulating Propionibacterium Breast cancer - CA Novel human CASB74 Human CASB7439 pro Propionibacterium Novel human CASB74 Propionibacterium

Heparin binding pe Heparin binding pe H. influenzae Hap Fragment of human Propionibacterium Novel human secret Human lung tumour L801P lung tumour Human lung cancer-

Title: Perfect score:

Sequence:

OM protein

Run on:

Scoring table:

Word size :

Searched:

Database :

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_'note= "This residue has a side chain
C(O)-NepsilonH-(CH2)3-Tris-ArgHel#3, where
the Tris-ArgHel#3 is represented in AAB71431"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide Arg Helix #3 for construction of Tris-Arg helix #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ALIGNMENTS
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ABP57745
                                                              AAU10338
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                                               ABJ10474
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                                                                                                                                                                                                                                                                                                                                                                                                                    AAU50234
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                                                                                                                               AAU51
                                                                                                                                                                                                         AAU61
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                                                                                                                                                                                                                                        AAB7
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Key
Modified-site
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Peptide Tris-Arg H
Heparin binding pe
Peptide Bis-Arg He
Heparin binding pe
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                                                                                                   August 9, 2003, 16:11:13; Search time 45.2571 Seconds (without alignments) 56.115 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | SIDSI/gcdata/geneseq/geneseqp-embl/Aa1980.DAT:*
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/SIDSI/gcgdata/geneseg/genesegp-emb1/AA2002.DAT:
/SIDSI/gcgdata/geneseg/genesegp-emb1/AA2003.DAT:
                                                                                                                                                                                                                                                                                                                                                1107863
        GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd
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                                                                                                                                                                                                                                                                                1107863 seqs, 158726573 residues
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                                                                        protein search, using sw model
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AAYB71428
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Gapop 60.0 , Gapext 60.0
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Result છું

Zea mays protein f Novel human diagno Fragment of human Human ORFX ORP2144 Propionibacterium

Lung cancer therap Human ORFX ORF2230

polype anti

Human DITHP po Human ovarian

C glutamicum prote A. thaliana EIN6 a Novel human diagno Amino acid sequenc Human protein sequ Human mitochondria A. thaliana EIN6 ( Rice AINTEGUMENTA-

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immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2, Immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2, Tetra-Arg Helix 2 for the manufacture of a medicament for the treatment of sepsia and the detection and removal of endotoxins. The peptides of the invention are used in a method for detecting endotoxin in a sample comprising contacting the sample with a labelled contotxin. The peptides can also be used in a method for removing endotoxin. The peptides can also be used in a method for removing endotoxin in a sample which comprises exposing the sample to a helix peptide, bound to a solid support, then collecting the sample to a helix peptide, bound to a solid support, then collecting the sample to a finity trap for endotoxins in e.g. dialysis-type treatments, or for affinity trap for endotoxins in e.g. dialysis-type treatments, or for affinity trap for endotoxins in e.g. dialysis-type treatments, or for affinity trap for endotoxins in e.g. dialysis-type treatments, or for a finite as model frameworks for endotoxin binding from which new analogues may be designed. This sequence represents the peptide Arg Helix #3 which is used in the construction of the invention.
                                                                                   treatment of sepsis and for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sepsis; branched chain peptide; antibacterial; immunosuppressive; endotoxin; helix peptide.
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16
/note= "Ala is modified by unidentified Rl group"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93.8%; Score 15; DB 23; L
100.0%; Pred. No. 8.1e-07;
Live 0; Mismatches 0;
                                                                            Use of cationic helix peptides for t
detection and removal of endotoxins
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                                                                                                                               Disclosure, Fig 2; 18pp; English.
                  Wolz G;
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Best Local Similarity 100.
Matches 15; Conservative
                 Wolz RL,
                                               WPI; 2002-659478/71.
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Modified-site
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Use of cationic helix peptides for treatment of sepsis and for the

(COMM-) COMMONWEALTH BIOTECHNOLOGIES INC

Harris RB, Wolz RL,

14-FEB-2001; 2001US-268410P.

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Into invention describes a novel use of antibacterial and immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2, Tetra-Arg Helix 3 for the manufacture of a medicament of Tetra-Arg Helix 3 for the manufacture of a medicament for the treatment of sepsis and the detection and removal of endotoxins. The peptides of the invention are used in a method for detecting endotoxin in a sample comprising contacting the sample with a labelled contacting the presence of any labelled molecule bound to endotoxin. The peptides can also be used in a method for removing endotoxin in a sample which comprises exposing the sample to a helix peptide. Dound to a solid support, then collecting the sample to a helix peptides may be in vivo, or the peptides may be used to form an affinity trap for endotoxins in e.g. dialysistype treatments, or for conved as model frameworks for endotoxin binding from which new analogues may be designed. This sequence represents the peptide Arg Helix #3 which is used in the construction of Tishs Helix #3, a branched chain
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bleeding wound; vascular anastomoses; leaking prosthetic vascular graft;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel heparin binding molecules (I). The molecules (I) are useful as heparin antagonist drugs for cardiovascular application and specifically neutralize heparin's conventional anticoagulant properties. (I) are also useful for counteracting actions of heparin locally e.g. in bleeding wounds, vascular anastomoses or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New heparin binding molecules, useful for reducing heparin content in a mammal by reducing the anticoagulant effects of heparin
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       detection and removal of endotoxins
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                                                                                                                                     Disclosure; Fig 1B; 18pp; English.
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immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2, immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2, retra-Arg Helix 2 or Tris-Arg Helix 3 for the manufacture of a medicament for the treatment of sepsis and the detection and removal of endotoxins. The peptides of the invention are used in a method for detecting endotoxin in a sample comprising contacting the sample with a labelled helix peptides and then detecting the presence of any labelled molecule bound to endotoxin. The peptides can also be used in a method for bound to endotoxin in a sample which comprises exposing the sample to a removing endotoxin in a sample which comprises exposing the sample to a endotoxin removal may be in vivo, or the peptides may be used to form an endotoxin removal may be in vivo, or the peptides may be used to form an affinity trap for endotoxins in e.g. dialysis-type treatments, or for removal of endotoxins from plasma fractionation products. They are also
pharmaceutical composition with insulin, as a substitute for protamine pharmaceutical composition with insulin, as a substitute for protamine for use in treating diabetics. The heparin binding molecules (1) specifically neutralize heparin's conventional anticoagulant properties without causing deleterious hemodynamic side-effects or exacerbation of the proliferative vascular response to injury. (1) are short-duration, intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations which intravenous drugs to be used in elective or emergency situations.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gapa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sepsis; branched chain peptide; antibacterial; immunosuppressive; endotoxin; helix peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 21; Length 19; 9.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.8%; Score 15; DB 100.0%; Pred. No. 9.7 1ve 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide Bis-Arg Helix #2 fragment #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 1A; 18pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB71428 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wolz G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 RRAARAARRARAEA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-NOV-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-659478/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 AB
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Wodifled-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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AMB
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This invention describes novel heparin binding molecules (I). The molecules (I) are useful as heparin antagonist drugs for cardiovascular conclections (I) are useful as heparin attagonist drugs for cardiovascular application and specifically neutralize heparin's conventional conference of heparin locally e.g. in bleeding wounds, vascular ansatomoses or conference of heparin locally e.g. in bleeding wounds, vascular ansatomoses or conference or conference of heparin composition with insulin, as a substitute for protamine of pharmaceutical composition with insulin, as a substitute for protamine of specifically neutralize heparin's conventional anticoagulant properties specifically neutralize heparin's conventional anticoagulant properties without causing deleterious hemodynamic side-effects or exacerbation of the proliferative vascular response to injury. (I) are short-duration, the proliferative vascular response to injury. (I) are short-duration, can safely and specifically neutralize heparin's proliferative response to injury. This sequence represents a heparin-binding peptide described in the method of the invention.
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used as model frameworks for endotoxin binding from which new analogues may be designed. This sequence represents the peptide Arg Helix #2 which is used in the construction of Bis-Arg Helix #2, a branched chain peptide described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Heparin binding peptide; antagonist; cardiovascular; coagulant; bleeding wound; vascular anastomoses; leaking prosthetic vascular graft; protamine substitute; treatment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New heparin binding molecules, useful for reducing heparin content in mammal by reducing the anticoagulant effects of heparin .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                       Length 19;
                                                                                                                                                         Indels
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100.0%; Pred. No. 0.15;
tive 0; Mismatches
                                                                                                                          DB 23; L 9.7e-07;
                                                                                                                                        100.0%; Pred. wc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (COMM-) COMMONWEALTH BIOTECHNOLOGIES INC.
                                                                                                                            Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                        Heparin binding peptide Arg helix #1.
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                                                                                                                                                                                                                                                                                                          AAY87834 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99EP-0119514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0166930
                                                                                                                               93.88;
                                                                                                                                                                                           2 RRAARAAARRARAEA 16
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                                                                                                                                                Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                 19 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Harris RB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                      Seguence
                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                  RESULT
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RESULT 6

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This invention describes novel recombinant antigens and their encoding nucleic acids derived from Mycobacterium tuberculosis. The novel polypeptides are useful for detecting M. tuberculosis infection in a biological sample by detecting antibodies which bind with the polypeptides, and are useful as vaccines for immunizing against M. tuberculosis infection. The new detection methods are needed as current vaccination strategies do not provide 100% immunity.
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                                                                               Antigen; diagnosis; detection; infection; antibody; immunisation; vaccine; immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New polypeptide comprising antigenic portions of M. tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Campos-Neto A, Dillon DC, Hendrickson RC, Houghton R;
Lodes MJ, Reed SG, Skeiky YAW, Twardzik DR, Vedvick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     M. tuberculosis and M. leprae marker protein #339.
                            M. tuberculosis recombinant antigen protein MO-4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 9; DB 2(; Pred. No. 0.5; 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU05688 standard; Protein; 105 AA.
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                                                                                                                                                                                                                                                                                                                                                           98US-0072596.
98US-0024753.
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                                                                                                                                                           Mycobacterium tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-APR-2003 (first entry)
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Mycobacterium leprae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Similarity 100...
5. Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 AAARRARAE 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                        (CORI-) CORIXA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VO200274903-A2.
                                                                                                                                                                                                                                                                                                            17-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                           05-MAY-1998;
18-FEB-1998;
                                                                                                                                                                                                        W09942118-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes polypeptides comprising an immunogenic part of a Mycobacterium tuberculosis antigen (Ag). Also described are vaccines and fusion protein containing M. tuberculosis Ag's.
M. tuberculosis Ag's. DNAs encoding them, derived fusion proteins and other polypeptides fragments, can be used in pharmaceutical compositions or vaccines to generate a protective or therapeutic immune response to M. tuberculosis and as respents in skin tests for diagnosis of tuberculosis. Ag can induce proliferation of, or cytokine secretion by, T. B or natural killer cells and/or macrophages in tuberculosis-immune subjects. AAZ19249 to AAZ19460 and AAY39083 to AAX39225 are used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antigens from Mycobacterium tuberculosis useful in diagnostic
skin tests and protective or therapeutic vaccines or compositions
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                                                                                                                                                                                                                                                                                                                                                                              s; M. tuberculosis; antigen; immun
immunisation; vaccine; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Campos-Neto A, Dillon DC, Hendrickson RC, Houghton R;
Lodes MJ, Reed SG, Skeiky YAW, Twardzik DR, Vedvick TS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 92;
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                                                                                                                                                                                                                                                                                                                         M. tuberculosis antigen 5' MO-4 amino acid sequence.
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5. 0.52;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5, Page 214; 299pp; English.
                                                                                                                                                                      AAX39179 standard; Protein; 92 AA.
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                                                                                                                                                                                                                                                                                                                                                                           Mycobacterium tuberculosis; M. immunotherapy; diagnosis; immu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               56.2%; S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-US03268
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98US-0025197
                                                                                                                                                                                                                                                                                                                                                                                                                                immune response; skin test.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mycobacterium tuberculosis
                                                                                                                                                                                                                                                                          (first entry)
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                          4 AARAAARRA 12
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4 AARAAARRA 12
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N-PSDB; AAZ19371.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORI-) CORIXA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W09942076-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-MAY-1998;
18-FEB-1998;
                                                                                                                                                                                                                                                                          05-NOV-1999
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                                                                                                                                                                                                                        AAY39179;
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Matches

AAY39036

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Gaps

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Length 92; 0; Indels ö

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RESULT 10
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                                                                                                                                                                             for survival or virulence of mycobacteria species. The method comprises aligning the genomic sequence of a first mycobacterium species on a genomic sequence of a second mycobacterium species and selecting a polynucleotide sequence that is highly conserved in both genomes with no counterparts in other bacterial genomic sequences and that corresponds to an essential gene for the survival or virulence of mycobacterium species. The method of the invention is useful for detecting M. tuberculosis or M. leprae infection. The method reduces the number of potential new targets and protective antigens for new drugs and vaccine of mompositions to treat and prevent mycobacterial diseases, particularly tuberculosis and leprosy. The present sequence represents a marker protein from Mycobacterium tuberculosis and Mycobacterium leprae identified using the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                 This invention relates to a novel method for identifying essential
                                                               Identifying and selecting genes for survival or virulence of mycobacteria by a comparative genomic analysis of the sequences of Mycobacterium tuberculosis and M. leprae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antigens from Mycobacterium tuberculosis useful in diagnostic skin tests and protective or therapeutic vaccines or compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mycobacterium tuberculosis; M. tuberculosis; antigen; immunogen; immunotherapy; diagnosis; immunisation; vaccine; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                    56.2%; Score 9; DB 23; Length 105; 100.0%; Pred. No. 0.57; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             M. tuberculosis antigen 5' MO-34 amino acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Campos-Neto A, Dillon DC, Hendrickson RC, Ho
Lodes MJ, Reed SG, Skeiky YAW, Twardzik DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 217-218; 299pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY39186 standard; Protein; 160 AA.
                                                                                                                                Claim 17; Page 536; 874pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US03268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0072967
98US-0025197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immune response; skin test.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mycobacterium tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 AAARRARAE 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19 AAARRARAE 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-527409/44.
N-PSDB; AAZ19378.
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                                                                                                                                                                                                                                                                                                                                                                                                                        105 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-MAY-1998;
18-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX39186;
Cole S;
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The present invention describes polypeptides comprising an immunogenic part of a Mycobacterium tuberculosis antigen (49). Also described are vaccines and fusion protein containing M. tuberculosis Ag's.

M. tuberculosis Ag's, DNAs encoding them, derived fusion proteins and other polypeptides fragments, can be used in pharmaceutical compositions or vaccines to generate a protective or therapeutic immune response to M. tuberculosis and as reagents in skin tests for diagnosis of tuberculosis. Ag can induce proliferation of, or cytokine secretion by, T. B or natural killer cells and/or macrophages in Ay39083 to tuberculosis-immune subjects. AAX19249 to AAX19460 and AAY39083 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes novel recombinant antigens and their encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acids derived from Mycobacterium tuberculosis. The novel polypeptides are useful for detecting M. tuberculosis infection in blological sample by detecting antibodies which bind with the polypeptides, and are useful as vaccines for immunizing against M. tuberculosis infection. The new detection methods are needed as
                                                                                                                                                                                                                                                                                     AAY39225 are used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antigen; diagnosis; detection; infection; antibody; immunisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polypeptide comprising antigenic portions of M. tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Houghton R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 160;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    M. tuberculosis recombinant antigen protein MO-34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Campos-Neto A, Dillon DC, Hendrickson RC, Ho
Lodes MJ, Reed SG, Skeiky YAW, Twardzik DR,
                                                                                                                                                                                                                                                                                                                                                                                           56.2%; Score 9; DB 20;
100.0%; Pred. No. 0.8;
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 9; DB 2C; Pred. No. 0.8; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Page 263; 323pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY39043 standard; Protein; 160 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 56.2%; Scur
100.0%; Pre
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mycobacterium tuberculosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                    Ouery Match
Best Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31 AAARRARAE 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 AAARRARAE 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-527416/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CORI-) CORIXA CORP.
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                            160 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vaccine; immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9942118-A2.
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18-FEB-1998;
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SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay;
                                                                                                                                                                                                                                                                                                                                                                                    Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful for treating acne vulgaris
                      Propionibacterium acnes immunogenic protein #7563.
                                                                                                                                                                                                                                                                                                           Skeiky YAW, Persing DH, Mitcham JL, Wang S.
L'maisonneuve J, Zhang Y, Jen S, Carter D;
                                                                                         dermatological; osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; SEQ ID No 7862; 1069pp; English.
                                                                                                                                                                                                                            21-APR-2000; 2000US-199047P.
02-JUN-2000; 2000US-208841P.
07-JUL-2000; 2000US-216747P.
                                                                                                                                                                                                  20-APR-2001; 2001WO-US12865.
27-FEB-2002 (first entry)
                                                                                                                        Propionibacterium acnes
                                                                                                                                                                                                                                                                                        (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-616774/71.
N-PSDB; AAS59534.
                                                                                                                                                WO200181581-A2.
                                                                                                                                                                          01-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a synthetic peptide which has a high affinity for glycominoglycans and proteoglycans. The peptide is useful in methods for modulating heparin or other glycominoglycans with anticoagulant activity, promoting call attachment or adhesion to natural or synthetic surfaces (especially vein grafts), modulating tumour cell metastasis, modulating cartilage differentiation, targeting drugs to epithelial cell surfaces (or to other cells expressing proteoglycans), modulating enzymes that act on glycoaminoglycan substrates, affinity, purification of bloactive sequences of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel synthetic peptides with high affinity for glycoaminoglycans and proteoglycans, useful for modulating heparin, promoting cell attachment, modulating tumour metastasis and modulating wound healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       glycominoglycan, modifying endothelial cell pro-coagulant or alti-coagulant functions mediated through glycominoglycans, and modulating wound healing. The peptide may also be used for blocking tissue uptake of heparin or other glycoaminoglycans in a mammal to increase heparin half-life in circulation.
                                                                                                                                                                                                        Glycoaminoglycan; proteoglycan; heparin modulation; anticoagulant; cell attachment; cell adhesion; vein graft; tumour cell metastasis; cartilage differentiation; wound healing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                 Peptide modulating activity of heparin, and other glycans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   50.0%; Score 8; DB 21; 100.0%; Pred. No. 0.91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      red. No. 0.9
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                   Schick BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 31; 76pp; English.
                                                                                                       AAB08179 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Verrecchio A,
                                                                                                                                                                                                                                                                                                                                                                                                         (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                 99US-0118276.
                                                                                                                                                                                                                                                                                                                                                     02-FEB-2000; 2000WO-US02853
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                                                                                                                                                               (first entry)
                           11111111
31 AAARRARAE 39
              7 AAARRARAE 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-543446/49.
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hes 8; Conserv
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                                                                                                                                                                                                                                                                                                   WO200045831-A1
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                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                     AAB08179;
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Matches
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Bhatia A;

Wang SS,

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Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medial calculation caused by the treatment, prevention and diagnosis of medial calculations caused by acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyellits), uveitis and endophthalmitis. Pustulosis, hypertosis and osteomyellits), uveitis and the central carvant. however it is particularly involved in the inflammatory lesions system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a blinding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The and determining the amount of bound protein in the sample. The captured in the amount of bound protein in the sample. The downregulate expression and activity of P. acnes polypeptides and downregulate expression and activity of P. acnes polypeptides and charmostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (ELISA).

Specification, but was obtained in electronic format directly from WIPO
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . 50.0%; Score 8; D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABJ10474 standard; Protein; 262 AA.
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Best Local Similarity 100.0%;
Matches 8; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             71 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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AAU46667 standard; Protein; 71 AA.

RESULT 12 AAU46667

AAU46667

7 AAARRARA 14 AAARRARA 8

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to the use of a human polynucleotide and polypeptide comprising at least 70 % identity to a 1791 base pair sequence over its entire length, or to a 193 residue amino acid sequence, both given in the specification, for the manufacture of a medicament for immunotherapeutically treating a patient suffering from or susceptible to breast carcinoma. The polynucleotides and polypeptides are useful as immunogens for specific prophylactic or therapeutic immunisation against breast tumours. The polynucleotide of the invention can be used in gene therapet 7 miss sequence represents a protein related to the CASB7439 breast cancer proteins of the invention.
                                                                                     Cytostatic; gene therapy; vaccine; immunotherapeutic; breast carcinoma; cancer; immunogen; immunisation; breast tumour; CASB7439.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Use of CASB7439 polypeptides and polynucleotides, their variants, immunogenic fragments and fusion proteins, for the manufacture of medicaments for the prophylaxis, treatment and diagnosis of breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 23; Length 262; 3. 8.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          colon cancer; tumour; immunoprophylaxis; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CASB7439; human; cytostatic; immunosuppressive; vaccine;
                                           Breast cancer - CASB7439 related protein SEQ ID No 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            colorectal cancer; immunogenic; autoimmune disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 8; I
                                                                                                                                                                                                                                                                                                                                                                               (SMIK ) SMITHKLINE BEECHAM BIOLOGICALS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 70; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAU10338 standard; Protein; 262 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human CASB7439 protein #3.
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                                                                                                                                                                                                                                                                                        15-FEB-2002; 2002WO-EP01649
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  21-NOV-2002 (first entry)
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Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tumors or cancer
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                                                                                                                                                                                                 WO200266506-A2.
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                                                                                                                                                       Homo saptens
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                                                                                                                                                                                                                                               29-AUG-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 14
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The invention relates to a novel isolated CASB7439 polypeptide (I).

(I), and its related polynuclectide (II) are useful in the manufacture of a vaccine for immunotherapeutically treating a patient suffering from or susceptible to carcinoma, preferably colon cancer or other coloncassociated tumours or diseases. (I), (II), and antibody to (I) are useful for the treatment of a subject by immunoprophylaxis or therapy by invitro induction of immune responses to (I), preferably for the treatment of colorectal cancer. An immunogenic composition composition comprising (I) is useful in medicine and for inducing an immune response or against human (ASB7439 polypeptide. (I) is useful as nimmunogen to produce antibodies immunospecific for (I), to identify membrane bound or soluble receptors, and in a method for the structure-based design of an colorectal and for inhibitor of (I), (I) or (II) can be used in a signification probps for conditions. (I) or (II) is useful for inducing, reconforting or modulating an immune response in a mammal. (II) is useful as conditional and genomic clones encoding (I). (II) its useful for staging concerous disorders and grading the nature of the cancerous tissue, and for thromosome localisation. Antibody to (I) is useful to isolate or identify clones expressing (I), to purify (I), and to prevent or treated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    identify clones expressing (I), to purify (I), and to prevent or treat cancer, particularly colorectal cancer, autoimmuse disease and related conditions. The present sequence represents the amino acid sequence of human CASB7439 polypeptide #3 as described in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                novel isolated CASB7439 polypeptide useful in diagnostics, and as vaccines for prophylactic and therapeutic treatment of cancers, particularly colorectal cancers, autoimmune diseases and related conditions
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                                                                                                                                                                                                                               Gaulis SRJ;
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100.0%; Pred. No. 8.6;
11ve 0; Mismatches
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                                                                                                                                                     (SMIK ) SMITHKLINE BEECHAM BIOLOGICALS.
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                                                                                                                                                                                                                               Cassart J,
23-FEB-2000; 2000GB-0004269.
20-APR-2000; 2000GB-0009905.
25-AUG-2000; 2000GB-0021080.
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Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                             WPI; 2002-041150/05.
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AAE33614
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WO200292627-A2.
                           21-NOV-2002.
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16-MAY-2001; 2001GB-0011974

07-MAY-2002; 2002WO-EP05011.

(GLAX ) GLAXOSMITHKLINE BIOLOGICALS SA.

Gaulis SRJ, Vinals De Bassols YC; Coche T,

WPI; 2003-120647/11. N-PSDB; AAD51534.

Use of a CASB7439 polynucleotide or polypeptide for manufacturing a medicament for immunotherapeutically preventing or treating a patient suffering from or susceptible to preneoplastic lesions of lung cancer and lung cancer.

Disclosure; Column 74-75; 55pp; English.

The invention relates to use of CASB7439 sequences for manufacturing a medicament for immunotherapeutically preventing or treating a patient suffering from or susceptible to preneoplastic lesions of lung cancer, and lung cancer and methods for diagnosing lesions of lung cancer are useful for manufacturing a medicament for treating preneoplastic lesions of lung cancer and lung cancer, such as SCLC, NSCLC (e.g. large cell (undifferentiated) carcinoma), squamous (spidermoid) carcinoma, carcinoma, carcinoma (including bronchoalveolar), bronchial gland tumours or mesorhellomas. CASB7439 DNA is used in gene therapy. The present sequence is human CASB7439 protein. 

262 AA; Sequence

Gaps 50.0%; Score 8; DB 24; Length 262; 100.0%; Pred. No. 8.6; 1.4e 0; Mismatches 0; Indels Query Match 50.0 Best Local Similarity 100. Matches 8; Conservative

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Search completed: August 9, 2003, 16:29:07 Job time : 45.2571 secs